Dotplot for stranded using anova_p_value HALLMARK_UNFOLDED_PROTEIN_RESPONSE HALLMARK_MTORC1_SIGNALING HALLMARK_MYOGENESIS HALLMARK_MYC_TARGETS_V2 HALLMARK KRAS SIGNALING UP HALLMARK G2M CHECKPOINT HALLMARK REACTIVE OXYGEN SPECIES PATHWAY HALLMARK_UV_RESPONSE_UP HALLMARK_E2F_TARGETS HALLMARK_INFLAMMATORY_RESPONSE HALLMARK MYC TARGETS V1 HALLMARK_TNFA_SIGNALING_VIA_NFKB HALLMARK_ESTROGEN_RESPONSE EARLY HALLMARK_ESTROGEN_RESPONSE_LATE HALLMARK COAGULATION HALLMARK_GLYCOLYSIS HALLMARK_IL2_STAT5_SIGNALING y value HALLMARK_XENOBIOTIC_METABOLISM HALLMARK_COMPLEMENT 0.2 HALLMARK_P53_PATHWAY 0.4 HALLMARK_APOPTOSIS 0.6 HALLMARK_TGF_BETA_SIGNALING HALLMARK APICAL JUNCTION 8.0 HALLMARK_CHOLESTEROL HOMEOSTASIS HALLMARK HYPOXIA HALLMARK PI3K AKT MTOR SIGNALING y_value HALLMARK ANGIOGENESIS HALLMARK EPITHELIAL MESENCHYMAL TRANSITION 8.0 HALLMARK HEME METABOLISM HALLMARK INTERFERON GAMMA RESPONSE 0.6 HALLMARK INTERFERON ALPHA RESPONSE 0.4 HALLMARK MITOTIC SPINDLE 0.2 HALLMARK_IL6_JAK_STAT3_SIGNALING HALLMARK UV RESPONSE DN HALLMARK ANDROGEN RESPONSE HALLMARK PANCREAS BETA CELLS HALLMARK HEDGEHOG SIGNALING HALLMARK ALLOGRAFT REJECTION HALLMARK NOTCH SIGNALING HALLMARK PROTEIN SECRETION HALLMARK KRAS SIGNALING DN HALLMARK WNT BETA CATENIN SIGNALING HALLMARK SPERMATOGENESIS HALLMARK DNA REPAIR HALLMARK APICAL SURFACE HALLMARK FATTY ACID METABOLISM HALLMARK ADIPOGENESIS HALLMARK PEROXISOME HALLMARK_BILE_ACID_METABOLISM · HALLMARK OXIDATIVE PHOSPHORYLATION 0.00 0.25 1.00 p_value